

OCT 0 9 2002

## **TECH CENTER 1600/2900**

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/043,539A

DATE: 09/23/2002

TIME: 10:23:43

Input Set : A:\EP.txt

Output Set: N:\CRF3\09232002\J043539A.raw

3 <110> APPLICANT: Cheung, Ambrose Manna, Adar Zhang, Gongyi 7 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR AFFECTING VIRULENCE DETERMINANTS IN BACTERIA 9 <130> FILE REFERENCE: DC-0199 11 <140> CURRENT APPLICATION NUMBER: US 10/043,539A C--> 12 <141> CURRENT FILING DATE: 2000-01-11 14 <150> PRIOR APPLICATION NUMBER: US 60/261,233 15 <151> PRIOR FILING DATE: 2001-01-12 17 <150> PRIOR APPLICATION NUMBER: US 60/261,607 18 <151> PRIOR FILING DATE: 2001-01-12 20 <150> PRIOR APPLICATION NUMBER: US 60/289,601 21 <151> PRIOR FILING DATE: 2001-05-08 23 <160> NUMBER OF SEQ ID NOS: 35 25 <170> SOFTWARE: PatentIn version 3.1 **ENTERED** 27 <210> SEQ ID NO: 1 28 <211> LENGTH: 947 29 <212> TYPE: DNA 30 <213> ORGANISM: Staphylococcus aureus 32 <220> FEATURE: 33 <221> NAME/KEY: CDS 34 <222> LOCATION: (208)..(552) 35 <223> OTHER INFORMATION: 38 <400> SEQUENCE: 1 39 gttttcaaaa tcggtggagg tgcatgaaaa agttattggg cattttttga aaataaaaaa 60 120 41 atatcaataa gttggagtca ttaccgaatt tttatactta tttgtttaga atgaacttta 43 taacatagtt ggatagagtt ttcgatttaa tacattaaat gtgaaccttg ctacaacaag 180 45 atgtgcatca gaaggagtgg tttaata atg agt aaa att aat gat att aat gat 234 Met Ser Lys Ile Asn Asp Ile Asn Asp 46 47

49 tta gtc aac gca aca ttt caa gtt aag aag ttt ttc aga gat aca aaa 50 Leu Val Asn Ala Thr Phe Gln Val Lys Lys Phe Phe Arg Asp Thr Lys

53 aag aag ttc aat ttg aac tat gaa gaa att tat att tta aat cat att 54 Lys Lys Phe Asn Leu Asn Tyr Glu Glu Ile Tyr Ile Leu Asn His Ile

57 tta aga agt gag tct aac gaa atc tca tct aaa gag att gct aag tgc 58 Leu Arg Ser Glu Ser Asn Glu Ile Ser Ser Lys Glu Ile Ala Lys Cys

61 tca gag ttc aaa cct tac tat tta act aaa gct tta caa aag cta aaa

62 Ser Glu Phe Lys Pro Tyr Tyr Leu Thr Lys Ala Leu Gln Lys Leu Lys 65 65 gat tta aaa ttg tta tca aag aaa aga agt tta caa gac gaa aga aca

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282

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66 Asp Leu Lys Leu Ser Lys Lys Arg Ser Leu Gln Asp Glu Arg Thr 67 75 80 85	
69 gtt att gtt tat gtt aca gat aca caa aaa gca aat att caa aaa ctg	522
70 Val Ile Val Tyr Val Thr Asp Thr Gln Lys Ala Asn Ile Gln Lys Leu	
71 90 95 100 105	
73 att toa gaa tta gaa gaa tao att aaa aat taaatcaagg ttaattgogt	572
74 Ile Ser Glu Leu Glu Glu Tyr Ile Lys Asn	
75 110 115	
77 ttaataacat tgaacgataa caatttatta atacgaagtt atttattcag cattgggaca	632
79 taaaattaac ttaaaattta aatattgaag atgctttaat taaagttaaa gaccagccat	692
81 accttatttc agcttattaa gcttgacaca aggtacacta gtctttttat tttaatattt	752
83 tottagaaaa toaagtttac gatcataaat attttctgcg atatagcttt ggatggttcc	812
85 aagtattttc tctataattt gtgtgcgata agcaaaaatt ctaactgcaa aaccatgtgt	872
87 aggcaattga gaaatagcaa cacgacaatc ggatgtattg ctataagaac taatggtttc	932
89 ataaactgaa tcgat	947
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94 <212> TYPE: PRT	
95 <213> ORGANISM: Staphylococcus aureus 97 <400> SEQUENCE: 2	
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100 I	
104 20 25 30	
107 Glu Glu Ile Tyr Ile Leu Asn His Ile Leu Arg Ser Glu Ser Asn Glu	
108 35 40 45	
111 Ile Ser Ser Lys Glu Ile Ala Lys Cys Ser Glu Phe Lys Pro Tyr Tyr	
112 50 55 60	
115 Leu Thr Lys Ala Leu Gln Lys Leu Lys Asp Leu Lys Leu Leu Ser Lys	
116 65 70 75 80	
119 Lys Arg Ser Leu Gln Asp Glu Arg Thr Val Ile Val Tyr Val Thr Asp	
120 85 90 95	
123 Thr Gln Lys Ala Asn Ile Gln Lys Leu Ile Ser Glu Leu Glu Glu Tyr	
124 100 105 110	
127 Ile Lys Asn	
128 115	
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133 <212> TYPE: PRT	
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138 Met Ala Ile Thr Lys Ile Asn Asp Cys Phe Glu Leu Leu Ser Met Val	
139 1 5 10 15 142 Thr Tyr Ala Asp Lys Leu Lys Ser Leu Ile Lys Lys Glu Phe Ser Ile	
142 Thr Tyr Ala Asp Lys Led Lys Sel Led Tie Lys Lys Gld File Sel Tie 143 20 25 30	
146 Ser Phe Glu Glu Phe Ala Val Leu Thr Tyr Ile Ser Glu Asn Lys Glu	
147 35 40 45	
150 Lys Glu Tyr Tyr Leu Lys Asp Ile Ile Asn His Leu Asn Tyr Lys Gln	
150 Lys old 1y1 1y1 Led Lys hisp 110 110 his 110 Led 110 110 110 110 110 110 110 110 110 11	
202	

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155		
158 159	Asp Lys Lys Arg Asn Glu His Asp Glu Arg Thr Val Leu Ile Leu Val 85 90 95	
162 163	Asn Ala Gln Gln Arg Lys Lys Ile Glu Ser Leu Leu Ser Arg Val Asn 100 105 110	
166 167	Lys Arg Ile Thr Glu Ala Asn Asn Glu Ile Glu Leu 115 120	
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	<211> LENGTH: 30	
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	<pre>&lt;220&gt; realore: &lt;223&gt; OTHER INFORMATION: Degenerate oligonucleotide.</pre>	
	<pre>&lt;223&gt; OTHER INFORMATION: Degenerate Offgondereotide: &lt;400&gt; SEQUENCE: 5</pre>	
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	<220> FEATURE:	
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	<212> TYPE: PRT <213> ORGANISM: Staphylococcus aureus	
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	<pre>&lt;222&gt; LOCATION: (1)(1)</pre>	
	<pre>&lt;222&gt; COTHER INFORMATION: "X" is defined as any amino acid residue.</pre>	
	<pre>&lt;220&gt; FEATURE:</pre>	

RAW SEQUENCE LISTING

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230 <221> NAME/KEY: MISC\_FEATURE 231 <222> LOCATION: (13)..(13) 232 <223> OTHER INFORMATION: "X" is defined as being either S or T. 235 <400> SEQUENCE: 8 🎶 237 Xaa Lys Ile Asn Asp Ile Asn Asp Leu Val Asn Ala Xaa Phe 241 <210> SEQ ID NO: 9 242 <211> LENGTH: 8 243 <212> TYPE: DNA 244 <213> ORGANISM: Staphylococcus aureus 246 <400> SEQUENCE: 9 247 aggagtgg 8 250 <210> SEQ ID NO: 10 251 <211> LENGTH: 6 252 <212> TYPE: DNA 253 <213> ORGANISM: Staphylococcus aureus 255 <400> SEQUENCE: 10 256 tagaat 6 259 <210> SEQ ID NO: 11 260 <211> LENGTH: 6 261 <212> TYPE: DNA 262 <213> ORGANISM: Staphylococcus aureus 264 <400> SEQUENCE: 11 265 ttaccg 6 268 <210> SEQ ID NO: 12 269 <211> LENGTH: 20 270 <212> TYPE: DNA 271 <213> ORGANISM: Staphylococcus aureus 273 <400> SEQUENCE: 12 274 ttactaaatt aaaaaaatta 20 277 <210> SEQ ID NO: 13 278 <211> LENGTH: 28 279 <212> TYPE: DNA 280 <213> ORGANISM: Staphylococcus aureus 282 <400> SEQUENCE: 13 283 taaattaatg ttatttttta ataattta 28 286 <210> SEQ ID NO: 14 287 <211> LENGTH: 8 288 <212> TYPE: DNA 289 <213> ORGANISM: Staphylococcus aureus 291 <400> SEQUENCE: 14 292 taaattaa 8 295 <210> SEQ ID NO: 15 296 <211> LENGTH: 8 297 <212> TYPE: DNA 298 <213> ORGANISM: Staphylococcus aureus 300 <400> SEQUENCE: 15

304 <210> SEQ ID NO: 16

301 ataattta

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· RAW SEQUENCE LISTING

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Input Set : A:\EP.txt

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(6.130 (6.	
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309 <400> SEQUENCE: 16	
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339 atatcaataa gttggagtca ttaccgaatt tttatactta tttgtttaga atgaacttta	120
341 taacatagtt ggatagagtt ttcgatttaa tacattaaat gtgaaccttg ctacaacaag	180
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27C (212) ADALYTON AL. 1 1	

378 <400> SEQUENCE: 23

376 <213> ORGANISM: Staphylococcus aureus

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/043,539A

DATE: 09/23/2002 TIME: 10:23:44

Input Set : A:\EP.txt

Output Set: N:\CRF3\09232002\J043539A.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the  $\langle 220 \rangle$  to  $\langle 223 \rangle$  fields of each sequence which presents at least one n or Xaa.

Seq#:8; Xaa Pos./1,13

 $\overline{\text{Seq}}$ :26; Xaa Pos. 2,4,10,15,16,17,19,20,24,25,31,38,44,45,47,51,52,58,59,61

Seq#:26; Xaa Pos. 73,77,78,86,87,95,97,99,102,106,110,114,117,118,120

## Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7